

# DEN\_WT\_BR3

%features

60

40

20

0

2.9

1.9

1

0

- % in genome
- detected
- % in sample

protein\_coding

pseudogene

lincRNA

antisense

miRNA

snoRNA

snRNA

IG\_V\_gene

processed\_transcript

misc\_RNA

rRNA

sense\_intronic

IG\_J\_gene

IG\_C\_gene

non\_coding

polymorphic\_pseudogene

sense\_overlapping

IG\_D\_gene

3prime\_overlapping\_ncrna

ncrna\_host

